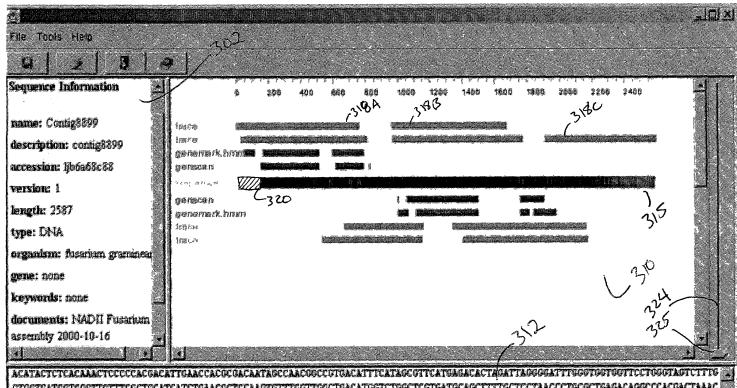


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F19.2



select	all			10 TO THE
Select:	Row name	accession	start	end description
D.	1 trace	<u> </u>	967	1775 sequence name fus_140_g24.b. 2a.t7a sequence start: 1 sequence end: 803 sequence
D E	2 trace	19699c98a	1339	2164 sequence name fus_141_g24.g 2a.sp6a sequence start. 2 sequence end: 819 sequence
O	3 trace	17ccb9464	668	1161 sequence name fus_143_j01.b. 2a:17a sequence start: 28 sequence end: 516 sequence
ON	4 trace	1048276b7	20	612 sequence name fus_143_j01.g. 2a spóa sequence start: 25 sequence end: 803 sequence
ΠŅ	5 trace	(b6d/shcad	1909	2599 sequence name fus_148_g13.b.2a.t7a sequence start. I sequence end: 691 sequence
gras, [,]	41.		***	4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4

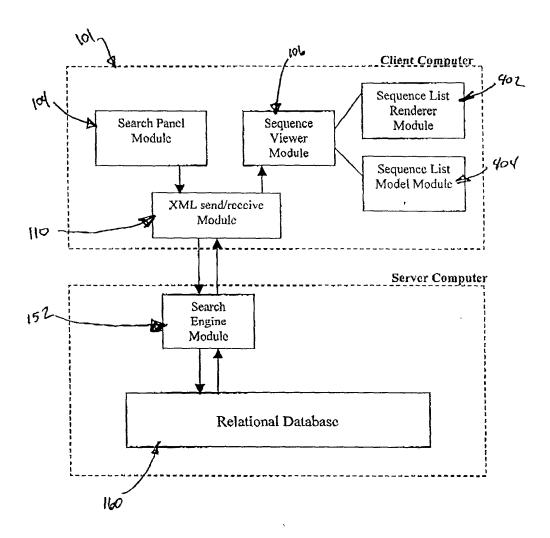
Welcome to the Feature Viewer

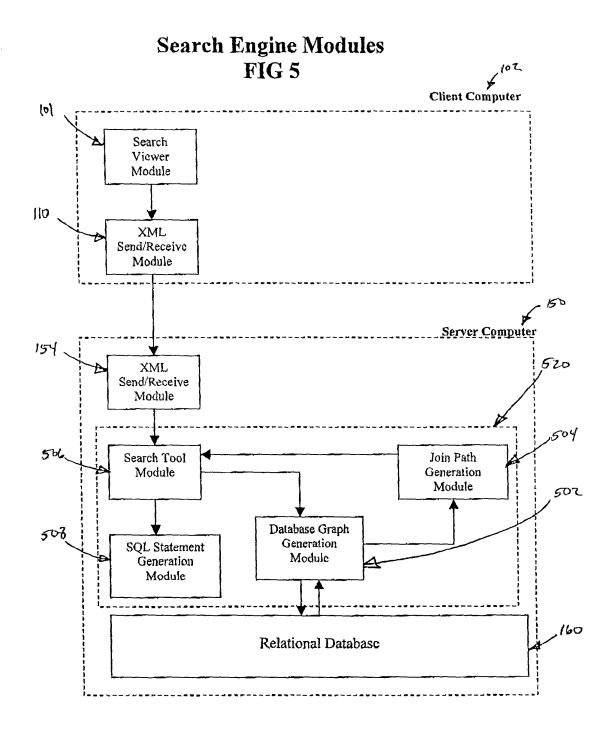
4

Fq. 3

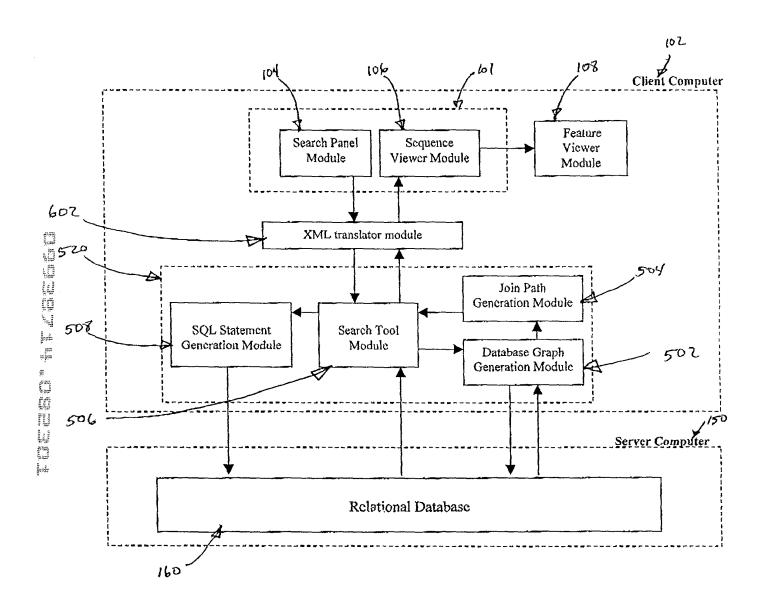
230

Search Viewer Modules FIG 4





Publicly Accessible Database Embodiment FIG 6



Process Flow Diagram

